

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: September 5, 2002, 13:01:12 : Search time 94.24 Seconds
(without alignments)
2687.268 Million cell updates/sec

Title: US-09-665-728-2

Perfect score: 1031

Sequence: 1 tctagcgacccttcggcc.....agcccgctcgagcgccgc 1031

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 383533 seqs, 122816752 residues

Total number of hits satisfying chosen parameters: 767066

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Issued_Patents_NA:*
1: /cgn2_6/pcdata/2/1na/5A_COMB.seq:*
2: /cgn2_6/pcdata/2/1na/5B_COMB.seq:*
3: /cgn2_6/pcdata/2/1na/5A_COMB.seq:*
4: /cgn2_6/pcdata/2/1na/5B_COMB.seq:*
5: /cgn2_6/pcdata/2/1na/PCUS_COMB.seq:*
6: /cgn2_6/pcdata/2/1na/backfile1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	50.4	4.9	2338	1	US-08-425-069-1
2	50.4	4.9	2338	2	US-08-317-844B-1
3	41.2	4.0	1185	3	US-09-023-339-3
4	41.2	4.0	1260	3	US-09-023-173-5
5	41.2	4.0	1308	3	US-09-023-173-10
6	41.2	4.0	1308	3	US-09-023-339-6
7	40.2	3.9	2214	3	US-08-864-038A-1
8	40.2	3.9	3331	3	US-08-864-038A-2
9	40.2	3.9	3331	3	US-08-864-038A-4
10	39.6	3.8	2502	1	US-08-073-384C-7
11	39.6	3.8	2502	1	US-08-254-359A-7
12	39.6	3.8	2502	1	US-08-483-043-7
13	39.6	3.8	2502	1	US-08-481-238-7
14	39.6	3.8	2502	2	US-08-471-066B-7
15	39.6	3.8	2502	2	US-08-484-956-7
16	39.6	3.8	2502	2	US-08-757-653-7
17	39.6	3.8	2502	2	US-08-599-491-7
18	39.6	3.8	2502	2	US-08-756-386-7
19	39.6	3.8	2502	2	US-08-823-516-7
20	39.6	3.8	2502	3	US-08-682-853A-7
21	39.6	3.8	2502	3	US-08-759-038-7
22	39.6	3.8	2502	3	US-08-758-314-7
23	39.6	3.8	2502	4	US-09-350-309-7
24	38.8	3.7	1929	4	US-09-380-420C-1
25	38.6	3.7	1598	1	US-08-211-682-24
26	38.4	3.7	2461	1	US-08-282-141-1
27	38.2	3.7	1995	1	US-08-425-069-3

28	38.2	3.7	1995	2	US-08-317-844B-3	Sequence 3, Appl1
29	37.6	3.6	71989	4	US-09-443-501A-2	Sequence 2, Appl1
30	37.4	3.6	6085	4	US-09-029-603-4	Sequence 4, Appl1
31	37.2	3.6	2260	2	US-08-788-750-1	Sequence 1, Appl1
32	37	3.6	2000	4	US-09-329-350-34	Sequence 34, Appl1
33	37	3.6	23673	4	US-09-773-816-1	Sequence 1, Appl1
34	36.8	3.5	50937	4	US-09-428-517-1	Sequence 10, Appl1
35	36.6	3.5	718	4	US-08-918-288-10	Sequence 10, Appl1
36	36.6	3.5	718	4	US-09-282-357-10	Sequence 10, Appl1
37	36.6	3.5	744	4	US-08-918-288-8	Sequence 8, Appl1
38	36.6	3.5	744	4	US-09-282-357-8	Sequence 8, Appl1
39	36.4	3.5	23673	4	US-09-773-816-1	Sequence 1, Appl1
40	36.2	3.5	2413	4	US-09-613-182-10	Sequence 10, Appl1
41	36.2	3.5	80161	3	US-09-036-987A-1	Sequence 1, Appl1
42	36.2	3.5	80161	4	US-09-370-700-1	Sequence 1, Appl1
43	36	3.5	432	1	US-08-642-255-48	Sequence 48, Appl1
44	36	3.5	756	1	US-08-642-255-50	Sequence 50, Appl1
45	36	3.5	2040	3	US-08-604-789B-1	Sequence 1, Appl1

ALIGNMENTS

RESULT 1
US-08-425-069-1
Sequence 1, 5728810
Patent No. 5728810
GENERAL INFORMATION:
APPLICANT: Lewis, Randolph V.
APPLICANT: Xu, Ming
APPLICANT: Himman, Michael B.
TITLE OF INVENTION: ISOLATED DNA CODING FOR SPIDER SILK
TITLE OF INVENTION: PROTEIN, A REPLICABLE VECTOR AND A TRANSFORMED CELL
TITLE OF INVENTION: CONTAINING THE ISOLATED DNA, AND PRODUCTS THEREOF
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolasch & Birch
STREET: 301 No. 5728810th Washington Street
City: Falls Church
STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22046
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/425,069
FILING DATE: 19-APR-1995
CLASSIFICATION: A35
ATTORNEY/AGENT INFORMATION:
NAME: Murphy Jr., Gerald M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1447-106P
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 205-8000
TELEFAX: (703) 205-8050
TELEX:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2338 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: Nephila clavipes
FEATURE:
NAME/KEY: CDS
LOCATION: 1..2154
OTHER INFORMATION: /product= "Nephila clavipes"

```

? OTHER INFORMATION: dragline silk protein"
?
? PUBLICATION INFORMATION:
?
? AUTHORS: Xu, Ming
?
? TITLE: Lewis, Randolph V.
?
? TITLE: Structure of a protein superfiber: Spider
?
? JOURNAL: Proc. Natl. Acad. Sci. U.S.A.
?
? VOLUME: 87
?
? PAGES: 7120-7124
?
? DATE: Sept.-1990
?
? RELEVANT RESIDUES IN SEQ ID NO: 1: FROM 1 TO 2338
?
? US-08-425-069-1

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Query Match	4.98;	Score 50.4;	DB 1;	Length 2338;
Best Local Similarity	49.38;	Pred. No. 0.0054;		
Matches 132; Conservative	0;	Mismatches 136;	Indels 0;	Gaps 0;

[illegible]

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Qy      754 atctgtctcagcagaccacgagagga 781
          | | | | |
Db      699 ACAAGCTGCTGACACAGAGCTGGAGCA 726

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RESULT 2

; Sequence 1, Application US/08317844B

GENERAL INFORMATION:

APPLICANT: Xu, Ming

TITLE OF INVENTION: ISOLATED DNA CODING FOR SPIDER SILK

TITLE OF INVENTION: PROTEIN, A REPLICABLE VECTOR AND A TRANSFORMED CELL
; TITLE OF INVENTION: CONTAINING THE ISOLATED DNA, AND PRODUCTS THEREOF
; TITLE OF INVENTION: CONTAINING THE ISOLATED DNA, AND PRODUCTS THEREOF

NUMBER OF SEQUENCES: 0.
CORRESPONDENCE ADDRESS:

ADDRESS: Blich, Stewart, Kolisch & Blich
STREET: 301 No. 5989894th Washington Street

CITY: Falls Church
STATE: Virginia

COUNTRY: U.S.A.
ZTP: 22046

```

; COMPUTER READABLE FORM:
; MEDTLM TYPE: F1000V

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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25
; C:\PROGRAM FILES\PATENTIN\DATA\

APPLICATION NUMBER: US/08/317,844B

CLASSIFICATION: 435

NAME: . Murphy Jr., Gerald M

REFERENCE/DOCKET NUMBER: 1447-105P

TELEPHONE: (703) 241-1300

TELEFAX: (703) 241-2848
TELEX: 248345
INFORMATION FOR SEQ ID NO.: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2338 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: *Nephila clavipes*
FEATURE:

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: LOCATION: 1. 2154
: OTHER INFORMATION: /product= "Nephila clavipes"
: OTHER INFORMATION: dragline silk protein"
: PUBLICATION INFORMATION:

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AUTHORS: Lewis, Randolph V.
TITLE: Structure of a protein superfiber: Spider
TITLE: draftline silk
JOURNAL: Proc. Natl. Acad. Sci. U.S.A.
VOLUME: 87

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;      DATE:  Sept.-1990
;      RELEVANT RESIDUES IN SEQ ID NO:  1: FROM 1 TO 2338
US-08-317-844B-1

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Query Match	4.9%;	Score 50.4;	DB 2;	Length 2338;
Best Local Similarity	49.3%;	Pred. No. 0.0054;		
Matches 132; Conservative	0;	Mismatches 136;	Indels 0;	Gaps 0;

QY	514	ttctggatgccccgtgcgaagactaccacaagaagaatgtgctctcctcgtcagctctctcagc	573
Db	459	tccttggaaagccaaagctgcagacacaggtggattatagtggaacaaagctccaaagctcagcagc	518
QY	574	tgaagcaatgcagatggtctctcgtctgtaacttggagactcagtgatgcagccagaccacccctg	633
Db	519	agcagcacccggagagtgctgtgacaaagcggatattacgggtggtcttgggtgacaaagtgccgg	578
QY	634	gtctacttggtcttggagagaatgcctccacacagttcttgccaagtgtccagcttgaca	693
Db	579	acaaagagagctatgagagacttggaaagccaaagctgctggacagagagatattagctgacaca	638
QY	694	agcagcagatgtcggagggtgttaagtctggttgagatccgcacatccagctccacaagtctc	753
Db	639	aggtgcagagctcagacagacagcagcagcagctggaggtggccgacaaagagagactaggtgc	698
QY	754	atctgctgtctcagacagacacagagaaga	781
Db	699	acaaagctcttgacacagacagcttgacga	726

RESULT

Sequence 3, Application US/09023339

GENERAL INFORMATION

APPLICANT: RODRIGUEZ, Raymond L.

TITLE OF INVENTION:	Production of 1-Antitrypsin
TITLE OF INVENTION:	in plants

; NUMBER OF SEQUENCES: 22
; CORRESPONDENCE ADDRESS:

ADDRESS: Dehlinger & Associates
STREET: P O Box 60850

CITY: Palo Alto
STATE: CA

COUNTRY: USA
ZTD: 04305

; COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

```

: MEDIUM TYPE: Diskette
: COMPUTER: IBM Compatible
: OPERATING SYSTEM: DOS
: SOFTWARE: FastSeq for Windows Version 2.0
: CURRENT APPLICATION DATA:
:   APPLICATION NUMBER: US/09/023,339
:   FILING DATE: 13-FEB-1998
:   PRIOR APPLICATION DATA:
:     APPLICATION NUMBER: 60/037,991
:     FILING DATE: 13-FEB-1997
:   ATTORNEY/AGENT INFORMATION:
:     NAME: Pelithory, Joanne R
:     REGISTRATION NUMBER: P42,995
:     REFERENCE/DOCKET NUMBER: 0665-0003.30
:   TELECOMMUNICATION INFORMATION:
:     TELEPHONE: 650-324-0880
:     TELEFAX: 650-324-0960
:   INFORMATION FOR SEQ ID NO: 3:
:     SEQUENCE CHARACTERISTICS:
:       LENGTH: 1185 base pairs
:       TYPE: nucleic acid
:       STRANDEDNESS: single
:       TOPOLOGY: linear
:   IMMEDIATE SOURCE:
:     CLONE: codon-optimized AAT coding sequence
:   US-09-023-339-3
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Query Match      4.0%; Score 41.2; DB 3; Length 1185;
Best Local Similarity 47.6%; Pred. No. 0.61;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;
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QY 464 catcaagctgcgcggtggtgccaacccgcgccttcacatcagctcttcggaagc 523
DB 664 CATCAGCAGCTGCAAGAAGCTCTCCAGCTGGGTGCTCTCATGAAGTACCTGGGAAGCC 743
QY 524 ccgtgccaaaggactaccacaagaagaatgagctcctgcagctcctgcagctgaagcagtc 583
DB 744 CACCGCATCTCTCTTCGCGGAGAGGAGGCAAGCTCCAGACCTCGAGAACGAGACTGAC 803
QY 584 gcaagtggtctcgctgctgactggtggaactcggtgacgcgacccaacctgctacctgac 643
DB 804 GCACGACATCATCAGCAAGATTCTCTGAGACGAGACAGGCGCTCCGCTAGCCTCCACCT 863
QY 644 tttagagagatgcgctccaccagcttcgccaagtgttcacagctcgggacaagcagcaggt 703
DB 864 CCCGAAGCTGAGCATACCGGCGACGTCAGACCTGAAGAGCGTGTGGCCAGCTGGGCAT 923
QY 704 gtcgagagtgttaa 717
DB 924 CACGAAGGCTCTCA 937
```

```

RESULT 4
US-09-023-173-5
: Sequence 5, Application US/09023173
: Patent No. 6066781
: GENERAL INFORMATION:
:   APPLICANT: Sulliff, Thomas D.
:   APPLICANT: Rodriguez, Raymond L.
:   TITLE OF INVENTION: Production of Mature Proteins
:   TITLE OF INVENTION: in Plants
:   NUMBER OF SEQUENCES: 23
:   CORRESPONDENCE ADDRESS:
:     ADDRESS: Dehlinger & Associates
:     STREET: 350 Cambridge Ave., Suite 250
:     CITY: Palo Alto
:     STATE: CA
:     COUNTRY: USA
:     ZIP: 94306
:   COMPUTER READABLE FORM:
:     MEDIUM TYPE: Diskette
:     COMPUTER: IBM Compatible
```

```

: OPERATING SYSTEM: DOS
: SOFTWARE: FastSeq for Windows Version 2.0
: CURRENT APPLICATION DATA:
:   APPLICATION NUMBER: US/09/023,173
:   FILING DATE: 13-FEB-1998
:   CLASSIFICATION: 435
:   PRIOR APPLICATION DATA:
:     APPLICATION NUMBER: 60/038,168
:     FILING DATE: 13-FEB-1997
:   ATTORNEY/AGENT INFORMATION:
:     NAME: Pelithory, Joanne R
:     REGISTRATION NUMBER: P42995
:     REFERENCE/DOCKET NUMBER: 0665-0007.30
:   TELECOMMUNICATION INFORMATION:
:     TELEPHONE: 650-324-0880
:     TELEFAX: 650-324-0960
:   INFORMATION FOR SEQ ID NO: 5:
:     SEQUENCE CHARACTERISTICS:
:       LENGTH: 1260 base pairs
:       TYPE: nucleic acid
:       STRANDEDNESS: single
:       TOPOLOGY: linear
:   IMMEDIATE SOURCE:
:     CLONE: codon-optimized Ramy3D signal-mature AAT
:   US-09-023-173-5
```

```

Query Match      4.0%; Score 41.2; DB 3; Length 1260;
Best Local Similarity 47.6%; Pred. No. 0.62;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;
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```

QY 464 catcaagctgcgcggtggtgccaacccgcgccttcacatcagctcttcggaagc 523
DB 739 CATCAGCAGCTGCAAGAAGCTCTCCAGCTGGGTGCTCTCATGAAGTACCTGGGAAGCC 818
QY 524 ccgtgccaaaggactaccacaagaagaatgagctcctgcagctcctgcagctgaagcagtc 583
DB 819 CACCGCATCTCTCTTCGCGGAGAGGAGGCAAGCTCCAGACCTCGAGAACGAGACTGAC 878
QY 584 gcaagtggtctcgctgctgactggtggaactcggtgacgcgacccaacctgctacctgac 643
DB 879 GCACGACATCATCAGCAAGATTCTCTGAGACGAGACAGGCGCTCCGCTAGCCTCCACCT 938
QY 644 tttagagagatgcgctccaccagcttcgccaagtgttcacagctcgggacaagcagcaggt 703
DB 939 CCCGAAGCTGAGCATACCGGCGACGTCAGACCTGAAGAGCGTGTGGCCAGCTGGGCAT 998
QY 704 gtcgagagtgttaa 717
DB 999 CACGAAGGCTCTCA 1012
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```

RESULT 5
US-09-023-173-10
: Sequence 10, Application US/09023173
: Patent No. 6066781
: GENERAL INFORMATION:
:   APPLICANT: Sulliff, Thomas D.
:   APPLICANT: Rodriguez, Raymond L.
:   TITLE OF INVENTION: Production of Mature Proteins
:   TITLE OF INVENTION: in Plants
:   NUMBER OF SEQUENCES: 23
:   CORRESPONDENCE ADDRESS:
:     ADDRESS: Dehlinger & Associates
:     STREET: 350 Cambridge Ave., Suite 250
:     CITY: Palo Alto
:     STATE: CA
:     COUNTRY: USA
:     ZIP: 94306
:   COMPUTER READABLE FORM:
:     MEDIUM TYPE: Diskette
:     COMPUTER: IBM Compatible
:     OPERATING SYSTEM: DOS
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SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,173
FILING DATE: 13-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/038,168
FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Petithory, Joanne R
REGISTRATION NUMBER: P42995
REFERENCE/DOCKET NUMBER: 0665-0007, 30
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-324-0880
TELEFAX: 650-324-0960
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 1308 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-09-023-173-10
```

```
Query Match 4.0%; Score 41.2; DB 3; Length 1308;
Best Local Similarity 47.6%; Pred. No. 0.63;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;
```

```
QY 464 catcaagcgcgcgtgaggtgtgcaaccccgctccttaatacagttcttcgagatgc 523
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 800 CATCACACACTGCAAGAGCTCTCCAGCTGGGTCTCTCATGAATCACTGGGGAACGC 859
QY 524 ccgtgccaagagactaccacaagaagaaatgagctcctgagctcctgagctgagcaagtc 583
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 860 CACCGCATCTCTTCTCTGCGGAGAGGCAAGCTCCAGACCTGAGAAAGAGCTGAC 919
QY 584 gcaagtgcttcgtctgactgagctgggagctcggtgacgcacccaccctgactcagc 643
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 920 GCACGACATCATCAAGAAATCTCTGGAAGAGAGGAGGCGCTCCGCTAGCTCCACT 979
QY 644 tttagagagatgcgcctccacagttcttgccaaagtgttccagctgagcaagcaagtc 703
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 980 CCCGAACTGAGCATCAACCGGACAGTACGACCTGAAGAGCTGTGGCCAGCTGGGCACT 1039
QY 704 gtccgagggtgttaa 717
    ||||| |||||
DB 1040 CACGAAGCTCTCA 1053
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```
RESULT 6
US-09-023-339-6
Sequence 6, Application US/09023339
Patent No. 6127145
GENERAL INFORMATION:
APPLICANT: Sulliff, Thomas D.
TITLE OF INVENTION: Rodriguez, Raymond L.
TITLE OF INVENTION: Production of '1-antitrypsin
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: P.O. Box 60850
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,339
```

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FILING DATE: 13-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/037,991
FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Petithory, Joanne R
REGISTRATION NUMBER: P42,995
REFERENCE/DOCKET NUMBER: 0665-0003, 30
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-324-0880
TELEFAX: 650-324-0960
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 1308 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: codon-optimized No. 61271451/XhoI fragment encoding Rany3D
US-09-023-339-6
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```
Query Match 4.0%; Score 41.2; DB 3; Length 1308;
Best Local Similarity 47.6%; Pred. No. 0.63;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;
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```
QY 464 catcaagcgcgcgtgaggtgtgcaaccccgctccttaatacagttcttcgagatgc 523
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 800 CATTCACACTGCAAGAGCTCTCCAGCTGGGTCTCTCATGAATCACTGGGGAACGC 859
QY 524 ccgtgccaagagactaccacaagaagaaatgagctcctgagctcctgagctgagcaagtc 583
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 860 CACCGCATCTCTTCTCTGCGGAGAGGCAAGCTCCAGACCTGAGAAAGAGCTGAC 919
QY 584 gcaagtgcttcgtctgactgagctgggagctcggtgacgcacccaccctgactcagc 643
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 920 GCACGACATCATCAAGAAATCTCTGGAAGAGAGGAGGCGCTCCGCTAGCTCCACT 979
QY 644 tttagagagatgcgcctccacagttcttgccaaagtgttccagctgagcaagcaagtc 703
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
DB 980 CCCGAACTGAGCATCAACCGGACAGTACGACCTGAAGAGCTGTGGCCAGCTGGGCACT 1039
QY 704 gtccgagggtgttaa 717
    ||||| |||||
DB 1040 CACGAAGCTCTCA 1053
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```
RESULT 7
US-08-864-038A-1
Sequence 1, Application US/08864038A
Patent No. 6001592
GENERAL INFORMATION:
APPLICANT: Kunio NAKASHIMA et al.
TITLE OF INVENTION: NOVEL POLYPEPTIDE GENE CDNA, VECTOR
TITLE OF INVENTION: CONTAINING SAID CDNA, HOST CELLS TRANSFORMED WITH SAID
TITLE OF INVENTION: VECTOR, POLYPEPTIDE PRODUCED THEREBY, METHOD OF PRODUCING
TITLE OF INVENTION: SAID POLYPEPTIDE, DNA ENCODING SAID POLYPEPTIDE AND ANTIBOD
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: 812-5 Hlrano
STREET: Isshinden
CITY: Tsu-city
STATE: Mie-prefecture
COUNTRY: JAPAN
ZIP: 514-01
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MB storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: Microsoft Windows 95
SOFTWARE: Word Perfect 6.1
CURRENT APPLICATION DATA:
```


TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-481-238-7

Query Match 3.8%; Score 39.6; DB 1; Length 2502;
Best Local Similarity 47.4%; Pred. No. 1.7;
Matches 108; Conservative 0; Mismatches 120; Indels 0; Gaps 0;

QY 179 gaagagcgtgtcccccacagagagacacccctgccttcctcagatgcacgg 238
DB 888 GGAGGCCCCCTGGCCCCCGCCGGAAGGGGCTTCGTGGGCTTGTCTCCCGCCGA 947
QY 239 ccccatgtggagcagatctgagtcaggtgacgtacagggcgccctcagcattctggagcgag 298
DB 948 GCCCATGTGGGCGGAGCTTCTGCCCCCTGGCCCCCGCCGGAAGGGGCGGCTCCACCGGGC 1007
QY 299 tctgagcagcgagcgaggtatcgcacactatgcgtgctgcttcacagaccaga 358
DB 1008 ACCAGACCCCTTTANGGGCTTANGGACCTNAGAGAGGTGGGGGNCCTCTCGCCAGGA 1067
QY 359 catggccagtgaccctcagcgagaccagtggtgttcagagaga 406
DB 1068 CCGGCCCTTTGGCCCTGAGGAGGAGGCTNAGACCTCMTGCCCCGGGGA 1115

RESULT 14
US-08-471-066B-7
Sequence 7, Application US/08471066B
Patent No. 5837450

GENERAL INFORMATION:

APPLICANT: Dahlberg, James E.
APPLICANT: Lyamichev, Victor I.
APPLICANT: Brow, Mary Ann D.
TITLE OF INVENTION: 5' Nucleases Derived From Thermostable
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Medlen & Carroll, LLP
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: California
COUNTRY: United States of America
ZIP: 94104

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/471.066B
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/254,359

FILING DATE: 06-JUN-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/073,384

FILING DATE: 04-JUN-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/986,330

FILING DATE: 07-DEC-1992

ATTORNEY/AGENT INFORMATION:
NAME: Ingolia, Diane E.
REGISTRATION NUMBER: 40,027

REFERENCE/DOCKET NUMBER: FORS-01800

TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
TELEFAX: (415) 397-8338

INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:

LENGTH: 2502 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-471-066B-7

Query Match 3.8%; Score 39.6; DB 2; Length 2502;
Best Local Similarity 47.4%; Pred. No. 1.7;
Matches 108; Conservative 0; Mismatches 120; Indels 0; Gaps 0;

QY 179 gaagagcgtgtcccccacagagagacacccctgccttcctcagatgcacgg 238
DB 888 GGAGGCCCCCTGGCCCCCGCCGGAAGGGGCTTCGTGGGCTTGTCTCCCGCCGA 947
QY 239 ccccatgtggagcagatctgagtcaggtgacgtacagggcgccctcagcattctggagcgag 298
DB 948 GCCCATGTGGGCGGAGCTTCTGCCCCCTGGCCCCCGCCGGAAGGGGCGGCTCCACCGGGC 1007
QY 299 tctgagcagcgagcgaggtatcgcacactatgcgtgctgcttcacagaccaga 358
DB 1008 ACCAGACCCCTTTANGGGCTTANGGACCTNAGAGAGGTGGGGGNCCTCTCGCCAGGA 1067
QY 359 catggccagtgaccctcagcgagaccagtggtgttcagagaga 406
DB 1068 CCGGCCCTTTGGCCCTGAGGAGGAGGCTNAGACCTCMTGCCCCGGGGA 1115

RESULT 15
US-08-484-956-7
Sequence 7, Application US/08484956
Patent No. 5843654

GENERAL INFORMATION:

APPLICANT: DAHLBERG, JAMES E.
APPLICANT: LYAMICHEV, VICTOR I.
APPLICANT: BROW, MARY ANN D.
APPLICANT: OLDENBURG, MARY C.
APPLICANT: HEISLER, LAURA
TITLE OF INVENTION: DETECTION OF p53 MUTATIONS
NUMBER OF SEQUENCES: 114
CORRESPONDENCE ADDRESS:
ADDRESSEE: HAVERSTOCK, MEDLEN & CARROLL
STREET: 220 MONTGOMERY STREET, SUITE 2200
CITY: SAN FRANCISCO
STATE: CALIFORNIA
COUNTRY: UNITED STATES OF AMERICA
ZIP: 94104

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,956
FILING DATE:
CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/402,601

FILING DATE: 09-MAR-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/337,164

FILING DATE: 09-NOV-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/254,359

FILING DATE: 06-JUN-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/073,384

FILING DATE: 04-JUN-1993

ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 07/986,330
FILING DATE: 07-DEC-1992
INFORMATION:

MOLECULE TYPE: DNA (genomic)
US-08-484-956-7

Query Match	3.88;	Score 39.6;	DB 2;	Length 2502;
Best Local Similarity	47.48;	Pred. No. 1.7;		
Matches 108; Conservative	0;	Mismatches 120;	Indels 0;	Gaps 0

OY	179	gaagaagcgtgccccgcacagaaagcaaaccccttgaccttcgtatgatcaag	238
Db	888	GGAGGCCCCCTGGCCCCCGGAAAGGGCCCTTTCGTGGGCTTTGTCCTTTCCCAGCCGA	947
OY	239	cctcacgtgagacatactgatacgagtgtgatcgacgycgactcaacgcatcttggagcgaag	298
Db	948	GCCCAATGTGGGCACGACTTCTGGCCCTTGCCGCCGCAAGGAGGGCCGGHTCACAGGGGC	1007
OY	299	tctgagcagcgcagaccggtcatacgccaatactgcgttgtcctttcaagaccaga	358
Db	1008	ACCAGA(C)CCTTTAANGGGCTTNAGGGACCTNNAAGGAGG(G)CGGGNC(N)CTGCGCAAGA	1067
OY	359	catltagccacagtaacctacaagggaaccagtgltgtttaagaaga	406
Db	1068	CCTGGCCCTTTTGGCCCTTGAGGAGGGCC(T)NACCTCNATGCCCGGGGA	1115

Search completed: September 5, 2002, 13:13:05
Job time: 713 sec

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: September 5, 2002, 13:13:05 : Search time 94.24 Seconds
(without alignments)
2150.336 Million cell updates/sec

Title: US-09-665-728-2_COPY_96_920
Perfect score: 825
Sequence: 1 atgacgcctagggcgcagct.....ctcctctctgatgcagac 825

Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0

Searched: 383533 seqs, 122816752 residues

Total number of hits satisfying chosen parameters: 767066

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	50.4	6.1	2338	1	US-08-425-069-1
2	50.4	6.1	2338	2	US-08-317-844B-1
3	41.2	5.0	1185	3	US-09-023-339-3
4	41.2	5.0	1260	3	US-09-023-173-5
5	41.2	5.0	1308	3	US-09-023-173-10
6	41.2	5.0	1308	3	US-09-023-339-6
7	40.2	4.9	2214	3	US-08-864-038A-1
8	40.2	4.9	3331	3	US-08-864-038A-2
9	40.2	4.9	3331	3	US-08-864-038A-4
10	39.6	4.8	2502	1	US-08-073-384C-7
11	39.6	4.8	2502	1	US-08-254-359A-7
12	39.6	4.8	2502	1	US-08-483-043-7
13	39.6	4.8	2502	1	US-08-481-238-7
14	39.6	4.8	2502	2	US-08-471-066B-7
15	39.6	4.8	2502	2	US-08-484-956-7
16	39.6	4.8	2502	2	US-08-757-653-7
17	39.6	4.8	2502	2	US-08-599-491-7
18	39.6	4.8	2502	2	US-08-756-386-7
19	39.6	4.8	2502	2	US-08-823-516-7
20	39.6	4.8	2502	3	US-08-682-853A-7
21	39.6	4.8	2502	3	US-08-759-038-7
22	39.6	4.8	2502	3	US-08-758-314-7
23	39.6	4.8	2502	4	US-09-350-309-7
24	38.6	4.7	1929	4	US-09-380-420C-1
25	38.6	4.7	1598	1	US-08-211-682-24
26	38.2	4.6	1995	1	US-08-425-069-3
27	38.2	4.6	1995	2	US-08-317-844B-3

28	37.4	4.5	6085	4	US-09-029-603-4	Sequence 4, Appl1
29	37	4.5	2000	4	US-09-329-350-34	Sequence 34, Appl1
30	36.8	4.5	50937	4	US-09-428-517-1	Sequence 1, Appl1
31	36.6	4.4	718	4	US-08-918-288-10	Sequence 10, Appl1
32	36.6	4.4	718	4	US-09-282-357-10	Sequence 10, Appl1
33	36.6	4.4	744	4	US-08-918-288-8	Sequence 8, Appl1
34	36.6	4.4	744	4	US-09-282-357-8	Sequence 8, Appl1
35	36.6	4.4	23673	4	US-09-773-816-1	Sequence 1, Appl1
36	36.2	4.4	2413	4	US-09-613-182-10	Sequence 10, Appl1
37	36.2	4.4	80161	3	US-09-036-987A-1	Sequence 1, Appl1
38	36.2	4.4	80161	4	US-09-370-700-1	Sequence 1, Appl1
39	36	4.4	432	1	US-08-642-255-48	Sequence 48, Appl1
40	36	4.4	756	1	US-08-642-255-50	Sequence 50, Appl1
41	36	4.4	2040	3	US-08-604-789B-1	Sequence 1, Appl1
42	36	4.4	2040	3	US-08-604-789B-12	Sequence 12, Appl1
43	36	4.4	2040	4	US-09-312-721A-1	Sequence 1, Appl1
44	36	4.4	2040	4	US-09-312-721A-12	Sequence 12, Appl1
45	35.4	4.3	4411529	4	US-09-103-840A-1	Sequence 1, Appl1

ALIGNMENTS

RESULT 1
US-08-425-069-1
; Sequence 1, Application US/08425069
; Patent No. 5728810
; GENERAL INFORMATION:
; APPLICANT: Lewis, Randolph V.
; APPLICANT: Xu, Ming
; APPLICANT: Hsman, Michael B.
; TITLE OF INVENTION: ISOLATED DNA CODING FOR SPIDER SILK
; TITLE OF INVENTION: PROTEIN, A REPLICABLE VECTOR AND A TRANSFORMED CELL
; TITLE OF INVENTION: CONTAINING THE ISOLATED DNA, AND PRODUCTS THEREOF
; NUMBER OF SEQUENCES: 69
; CORRESPONDENCE ADDRESSES:
; ADDRESS: Birch, Stewart, Kolasch & Birch
; STREET: 301 No. 5728810th Washington Street
; CITY: Falls Church
; STATE: Virginia
; COUNTRY: U.S.A.
; ZIP: 22046
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: Patent Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/425,069
; FILING DATE: 19-APR-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Murphy Jr., Gerald M
; REGISTRATION NUMBER: 28,977
; REFERENCE/DOCKET NUMBER: 1447-106P
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 205-8000
; TELEFAX: (703) 205-8050
; TELEX:
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2338 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; ORGANISM: Nephilia clavipes
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..2154
; OTHER INFORMATION: /product= "Nephilia clavipes"

? OTHER INFORMATION: dragline silk protein"
 ? PUBLICATION INFORMATION:
 ? AUTHORS: Xu, Ming
 ? TITLE: Structure of a protein superfiber: Spider
 ? TITLE: dragline silk
 ? JOURNAL: Proc. Natl. Acad. Sci. U.S.A.
 ? VOLUME: 87
 ? PAGES: 7120-7124
 ? DATE: Sept.-1990
 ? RELEVANT RESIDUES IN SEQ ID NO: 1: FROM 1 TO 2338
 US-08-425-069-1

Query Match	6.18;	Score 50.4;	DB 1;	Length 2338;
Best Local Similarity	40.39;	Prod NO	0.004;	

Matches 132; Conservative 0; Mismatches 136; Indels 0; Gaps 0;

[illegible]

RESULT 2
US-08-317-844B-1
; Sequence 1, Application US/08317844B
Date: 11/11/2008

PATENT NO. 5989894
GENERAL INFORMATION:
APPLICANT: Lewis, Randolph V.
APPLICANT: Xu, Ming
APPLICANT: Himan, Michael B.
TITLE OF INVENTION: ISOLATED DNA CODING FOR SPIDER SILK
TITLE OF INVENTION: PROTEIN, A REPLICABLE VECTOR AND A TRANSFORMED CELL
TITLE OF INVENTION: CONTAINING THE ISOLATED DNA, AND PRODUCTS THEREOF
NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSEE: Birch, Stewart, Kolach & Birch
STREET: 301 NO. 5989894th Washington Street
CITY: Falls Church
STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22046
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/317,844B
FILING DATE: 04-OCT-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Murphy Jr., Gerald M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1447-105P
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 241-1300

```

1 TELEFAX: (703) 241-2848
2
3 TELEX: 248345
4
5 INFORMATION FOR SEQ ID NO: 1:
6
7 SEQUENCE CHARACTERISTICS:
8     LENGTH: 2338 base pairs
9     TYPE: nucleic acid
10    STRANDEDNESS: single
11    TOPOLOGY: linear
12
13 MOLECULE TYPE: cDNA
14
15 HYPOTHETICAL: NO
16
17 ORIGINAL SOURCE:
18
19 ORGANISM: Nephilia clavipes
20
21 FEATURE:
22
23     NAME/KEY: CDS
24     LOCATION: 1..2154
25
26 OTHER INFORMATION: /product="Nephilia clavipes
27                   dragline silk protein"
28
29 PUBLICATION INFORMATION:
30
31     AUTHORS: Xu, Ming
32
33     TITLE: Structure of a protein superfiber: Spider
34
35     TITLE: dragline silk
36
37     JOURNAL: Proc. Natl. Acad. Sci. U.S.A.
38
39     VOLUME: 87
40
41     PAGES: 7120-7124
42
43     DATE: Sept.-1990
44
45     RELEVANT RESIDUES IN SEQ ID NO: 1: FROM 1 TO 2338
46
47 US-08-317-844B-1

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Query Match	6.18;	Score	50.4;	DB	2;	Length	2338;
Best Local Similarity	49.38;	Prod	NO	0	000.		

base local similarity 4.50, rec. no. 0.004,
Matches 132; Conservative 0; Mismatches 136; Indels 0; Gaps 0;

Accession	Sequence	Position
QY	ctgaagcaatcgcagaatggtctctcgtgctgaactgggagactggtgtaacgcagcccaaccctg	538
Db		
QY	agcacacaccccgagagctgctgcacacagcgcgatgacgggtgttggtagacaaagctccgg	578
Db		
QY	gctacctgaccttctgagagaatcgccctccacaagcttcctgaccaagtcttccagcttgaca	598
Db		
QY	acaaagagacgtatgagacacttggaaagccaaagcgtcgtggaagagaaagatttagcttgaca	638
Db		
QY	agcagcaaggtgtcggagagtgttaaatgttgtagtccgcacatcccaatccaggtccctcaaatc	658
Db		
QY	aggtgcacaggtccacacagcagcagcagcagctgcgaggtgcccgacaaagagagactgactgg	698
Db		
QY	atctgctgtctcagcagaccacgaagaaga	686
Db		
QY	acaaaggtctctgacacaaagcgtggaaca	726
Db		

RESULT 3
ME-00-022-220-2

US 02-023-535-5
Sequence 3, Application US/09023339
Patent No. 6127145
GENERAL INFORMATION:
APPLICANT: Sutliff, Thomas D.
APPLICANT: Rodriguez, Raymond L.
TITLE OF INVENTION: Production of '1-Antitrypsin
TITLE OF INVENTION: In Plants
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: P.O. Box 60850
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,339
FILING DATE: 13-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/037,991
FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Pelichory, Joanne R
REGISTRATION NUMBER: P42,995
REFERENCE/DOCKET NUMBER: 0665-0003.30
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-324-0880
TELEFAX: 650-324-0960
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 1185 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: codon-optimized AAT coding sequence
US-09-023-339-3

Query Match 5.0%; Score 41.2; DB 3; Length 1185;
Best Local Similarity 47.6%; Pred. No. 0.47;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;

QY 369 catcaagctgcctgaggttgccaaccccgctccctatctatcgtcttcgagatgc 428
DB 684 CATCCAGCACTGCAGAAAGTCTCCAGCTGGGTCTCTCATGAGTACTGGGGAACGC 743
QY 429 ccgtgccaagagctaccacaagaagatgagctcctgcaagctctcagctgagcaatgc 488
DB 744 CACCGCATCTTCTCTCGCGGACGAGGCAAGCTCCACACTGGAGAACAGAGCTGAC 803
QY 489 gcaagtgctctgctgactgagtgagctgagctgagctgagctgagctgagctgagctgagc 548
DB 804 GCACGACATCATCAGCAAGTCTCGGAGAGACGAGACGAGCGCTCCGCTCCACCT 863
QY 549 ttgttgagagatgctccaccagttctgccaagtgcttccagctgagcaagagcaggt 608
DB 864 CCCGAAGCTGAGCATCACCGGACGATGACGCTGAAGAGCGTGGCCAGCTGGGCGAT 923
QY 609 gtcgaggtgttaa 622
DB 924 CACGAAGCTCTTCA 937

RESULT 4
US-09-023-173-5
Sequence 5, Application US/09023173
Patent No. 6066781
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Thomas D.
APPLICANT: Rodriguez, Raymond L.
TITLE OF INVENTION: Production of Mature Proteins
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite 250
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,173
FILING DATE: 13-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/038,168
FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Pelichory, Joanne R
REGISTRATION NUMBER: P42995
REFERENCE/DOCKET NUMBER: 0665-0007.30
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-324-0880
TELEFAX: 650-324-0960
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1260 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: codon-optimized Ramy3d signal-mature AAT
US-09-023-173-5

Query Match 5.0%; Score 41.2; DB 3; Length 1260;
Best Local Similarity 47.6%; Pred. No. 0.48;
Matches 121; Conservative 0; Mismatches 133; Indels 0; Gaps 0;

QY 369 catcaagctgcctgaggttgccaaccccgctccctatctatcgtcttcgagatgc 428
DB 759 CATCCAGCACTGCAGAAAGTCTCCAGCTGGGTCTCTCATGAGTACTGGGGAACGC 818
QY 429 ccgtgccaagagctaccacaagaagatgagctcctgcaagctctcagctgagcaatgc 488
DB 819 CACCGCATCTTCTCTCGCGGACGAGGCAAGCTCCACACTGGAGAACAGAGCTGAC 878
QY 489 gcaagtgctctgctgactgagtgagctgagctgagctgagctgagctgagctgagctgagc 548
DB 879 GCACGACATCATCAGCAAGTCTCGGAGAGACGAGACGAGCGCTCCGCTCCACCT 938
QY 549 ttgttgagagatgctccaccagttctgccaagtgcttccagctgagcaagagcaggt 608
DB 939 CCCGAAGCTGAGCATCACCGGACGATGACGCTGAAGAGCGTGGCCAGCTGGGCGAT 998
QY 609 gtcgaggtgttaa 622
DB 999 CACGAAGCTCTTCA 1012

RESULT 5
US-09-023-173-10
Sequence 10, Application US/09023173
Patent No. 6066781
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Thomas D.
APPLICANT: Rodriguez, Raymond L.
TITLE OF INVENTION: Production of Mature Proteins
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite 250
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS

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SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023, 173
FILING DATE: 13-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/038, 168
FILING DATE: 13-FEB-1997
ATTORNEY/AGENT INFORMATION:
NAME: Petichory, Joanne R
REGISTRATION NUMBER: P42995
REFERENCE/DOCKET NUMBER: 0665-0007.30
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-324-0880
TELEFAX: 650-324-0960
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 1308 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
OS-09-023-173-10

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Query Match	5.0%	Score 41.2	DB 3	Length 1308
Best Local Similarity	47.6%	Pred. No. 0.48	Mismatches 133	Indels 0
Matches 121	Conservative 0	Mismatches 133	Indels 0	Gaps 0
QY	369	catcaagctccgctgtagaggtgcccaccccgctctctcatctacgtctctcgatgc	428	
Db	800	CATCCAGCAGCTGCAGGAAGCTCTCCAGCTGGTGCTCCCAAGAACTCTGGGAGAC	859	
QY	429	ccgtcccaagagactaccacaagaagaagtagtctctgcagctctcgcagctgaagcagc	488	
Db	860	CACGCCATCTTCTTCTCCCTGCCGAGAGAGGGCAAGCTCCAGCACTGGAGAACAGAGCTGAC	919	
QY	489	gcagagtggtcttcgtgtgctgtaactgaggactcggtgacgacaccacccctgtaacctgac	548	
Db	920	GCACGACATCATCAACGAAGTCTCTGGACAACAGAGACAGGCCCTCCGCAAGCTCCACCT	979	
QY	549	tttctgagagatacgctccacccagctctgcgcgaagtgttccacgtgagacacgaagtc	608	
Db	980	CCCGAAGCTGAGCATTCACACCGGCACCTACGACTGAAAGACGCTGCGCACACTGGGCAT	1039	
QY	609	gtcggagagtgcttaa	622	
Db	1040	CACGAAGGCTTTCA	1053	

RESULT 6
US-09-023-339-6
Sequence 6, Application US/09023339
Patent No. 6127145
GENERAL INFORMATION:
APPLICANT: Sutcliffe, Thomas D.
APPLICANT: Rodriguez, Raymond L.
TITLE OF INVENTION: Production of '1-Antitrypsin
TITLE OF INVENTION: In Plants
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: P.O. Box 60850
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023.339

```

?      FILING DATE: 13-FEB-1998
?      PRIOR APPLICATION DATA:
?      APPLICATION NUMBER: 60/037,991
?      FILING DATE: 13-FEB-1997
?      ATTORNEY/AGENT INFORMATION:
?      NAME: Petlinhory, Joanne R
?      REGISTRATION NUMBER: P42,995
?      REFERENCE/DOCKET NUMBER: 0665-0003.30
?      TELECOMMUNICATION INFORMATION:
?      TELEPHONE: 650-324-0980
?      TELEFAX: 650-324-0960
?      INFORMATION FOR SEQ ID NO: 6:
?      SEQUENCE CHARACTERISTICS:
?      LENGTH: 1308 base pairs
?      TYPE: nucleic acid
?      STRANDEDNESS: single
?      TOPOLOGY: linear
?      IMMEDIATE SOURCE:
?      CLONE: codon-optimized No. 612/71451/XhoI fragment encoding Rany2D
?      CLONE: signal peptide-AAT fusion protein
?      IS-09-023-339-6

```

Query Match	5.0%	Score 41.2	DB 3	Length 1308
Best Local Similarity	47.6%	Pred. No. 0.48		
Matches 121	Conservative	0	Mismatches 133	Indels 0
QY	369	catcaagagctgcgcgfygagagtggtcccaaccocgcgcctctcatcagtcgtcttcgcatgc	428	
Db	800	catccacgacactgcgaamaaagctctccagctggggtgcctcttgaaagacctggggaaacgc	859	
QY	429	ccgtgcgaagagactaccacaagaagaatgagctccctgcagctccctgcagctgaagcagtc	488	
Db	860	caccgcacatctcttcttcctcccgagcagagggcgaagctccagactggaaacagcagctgac	919	
QY	489	gagagtggtgttcgcgtgctgacttggagacttggtgagccgcaacccttgcttaactgac	548	
Db	920	gcacgcacatctatcagaaattccttgaaacagagacacggcgctccgctgacctccacct	979	
QY	549	ttttggagagatcgctccacacagcttcgcgcgaagtgttcacagcttgacaagcagcagtc	608	
Db	980	cccgaaagctgacacatcacccgacagctacgacctgaaagagcgtgctggccagactggggcatt	1039	
QY	609	gtcggaggtgttaa	622	
Db	1040	cacgaagctcttca	1053	

```

RESULT 7
US-08-864-038A-1
: Sequence 1, Application US/08864038A
: Patent No. 6001592
: GENERAL INFORMATION:
: APPLICANT: Kunio NAKASHIMA et al.
: TITLE OF INVENTION: NOVEL POLYPEPTIDE GENE CDNA, VECTOR
: TITLE OF INVENTION: CONTAINING SAID CDNA, HOST CELLS TRANSFORMED WITH SAID
: TITLE OF INVENTION: VECTOR, POLYPEPTIDE PRODUCED THEREBY, METHOD OF PRODUCING
: TITLE OF INVENTION: SAID POLYPEPTIDE, DNA ENCODING SAID POLYPEPTIDE AND ANTIBOD
: NUMBER OF SEQUENCES: 4
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: 812-5 Hirano
: STREET: Isshinden
: CITY: Tsu-city
: STATE: Mie-prefecture
: COUNTRY: JAPAN
: ZIP: 514-01
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MB storage
: COMPUTER: IBM Compatible
: OPERATING SYSTEM: Microsoft Windows 95
: SOFTWARE: Word Perfect 6.1
: CURRENT APPLICATION DATA:

```



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1      FILING DATE: 04-JUN-1993
2      CLASSIFICATION: 536
3      PRIOR APPLICATION DATA:
4      APPLICATION NUMBER: US 07/986,330
5      FILING DATE: 07-DEC-1992
6      ATTORNEY/AGENT INFORMATION:
7      NAME: Carroll, Peter G.
8      REGISTRATION NUMBER: 32,837
9      REFERENCE/DOCKET NUMBER: FORS-00613
10     TELECOMMUNICATION INFORMATION:
11     TELEPHONE: 415/705-8410
12     TELEFAX: 415/397-8338
13     INFORMATION FOR SEQ ID NO: 7:
14     SEQUENCE CHARACTERISTICS:
15     LENGTH: 2502 base pairs
16     TYPE: nucleic acid
17     STRANDEDNESS: single
18     TOPOLOGY: linear
19     US-08-073-384C-7

```

Query Match	4.8%;	Score 39.6;	DB 1;	length 2502;
Best Local Similarity	47.4%;	Pred. No. 1.3;		
Matches 108;	Conservative	0;	Mismatches 120;	Indels 0;
			Gaps	0;

[illegible]

RESULT 11
 US-08-254-359A-7
 Sequence 7, Application US/08254359A
 Patent No. 5614402
 GENERAL INFORMATION:
 APPLICANT: DAHLBERG, JAMES E.
 APPLICANT: LYAMICHEV, VICTOR I.
 APPLICANT: BROW, MARY ANN D.
 TITLE OF INVENTION: 5' NUCLEASES DERIVED FROM THERMOSTABLE
 TITLE OF INVENTION: DNA POLYMERASE
 NUMBER OF SEQUENCES: 40
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: HAVERTOCK, MEDLEN & CARROLL
 STREET: 220 MONTGOMERY STREET, SUITE 2200
 CITY: SAN FRANCISCO
 STATE: CALIFORNIA
 COUNTRY: UNITED STATES OF AMERICA
 ZIP: 94104
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/254,359A
 FILING DATE:
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/073,384
 FILING DATE: 06-JUN-1993
 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/986,330
FILING DATE: 07-DEC-1992
ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FORS-01000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 2502 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-254-359A-7

Query Match 4.8%, Score 39.6; DB 1; Length 2502;
Best Local Similarity 47.4%; Pred. No. 1.3;
Matches 108; Conservative 0; Mismatches 120; Indels 0; Gaps 0;

QY 84 gaagacgctgtcccccgcacagagagacgacccctgctcgtcttcgattgacagg 143
DB 888 GGAGGCCCCCTGGCCCCCGCGAAGGGGCTTCGTGGGCTTTGCTTCCCGCCCGA 947
QY 144 ctccatgtggagacatctgtagcaggtagtcgaagcgccctcaacgcatctggagcgag 203
DB 948 GCCCATGTGGGGCGAGCTTCTGGCCCTGCGCCGCGCAGGAGGGCCGGGTCCACCGGGC 1007
QY 204 tctgagcagcgacgacgggctgcatcgcacactatgctgtgcttcacagaccaga 263
DB 1008 ACCAGACCCCTTTANGGCCCTNAGGACCTNAGAGAGGTGCGGGNCTCTGCCAAGA 1067
QY 264 catggccagtgacacctcaacgagcgaccagtggtgttcagagaga 311
DB 1068 CCTGGCCGTTTGGCCCTGAGGAGGGCCCTNAGACCTCTGCGCCGGGGA 1115

RESULT 12
US-08-483-043-7
Sequence 7, Application US/08483043
Patent No. 5691142
GENERAL INFORMATION:
APPLICANT: Dahlberg, James E.
APPLICANT: Lyamichev, Victor I.
TITLE OF INVENTION: SYNTHESIS-DEFICIENT THERMOSTABLE DNA
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: HAVERSTOCK, MEDLEN & CARROLL
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: California
COUNTRY: United States of America
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,043
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/073,384
FILING DATE: 04-JUN-1993
APPLICATION NUMBER: US 07/986,330
FILING DATE: 07-DEC-1992
ATTORNEY/AGENT INFORMATION:

NAME: Carroll, Peter G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FORS-00613
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/705-8410
TELEFAX: 415/397-8338
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 2502 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-483-043-7

Query Match 4.8%, Score 39.6; DB 1; Length 2502;
Best Local Similarity 47.4%; Pred. No. 1.3;
Matches 108; Conservative 0; Mismatches 120; Indels 0; Gaps 0;

QY 84 gaagacgctgtcccccgcacagagagacgacccctgctcgtcttcgattgacagg 143
DB 888 GGAGGCCCCCTGGCCCCCGCGAAGGGGCTTCGTGGGCTTTGCTTCCCGCCCGA 947
QY 144 ctccatgtggagacatctgtagcaggtagtcgaagcgccctcaacgcatctggagcgag 203
DB 948 GCCCATGTGGGGCGAGCTTCTGGCCCTGCGCCGCGCAGGAGGGCCGGGTCCACCGGGC 1007
QY 204 tctgagcagcgacgacgggctgcatcgcacactatgctgtgcttcacagaccaga 263
DB 1008 ACCAGACCCCTTTANGGCCCTNAGGACCTNAGAGAGGTGCGGGNCTCTGCCAAGA 1067
QY 264 catggccagtgacacctcaacgagcgaccagtggtgttcagagaga 311
DB 1068 CCTGGCCGTTTGGCCCTGAGGAGGGCCCTNAGACCTCTGCGCCGGGGA 1115

RESULT 13
US-08-481-238-7
Sequence 7, Application US/08481238
Patent No. 5795763
GENERAL INFORMATION:
APPLICANT: DAHLBERG, JAMES E.
APPLICANT: LYAMICHEV, VICTOR I.
APPLICANT: BROW, MARY ANN D.
TITLE OF INVENTION: SYNTHESIS-DEFICIENT THERMOSTABLE DNA
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL
STREET: 220 MONTGOMERY STREET, SUITE 2200
CITY: SAN FRANCISCO
STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/481,238
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FORS-01798
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 2502 base pairs

